

228844066.ST25.txt  
SEQUENCE LISTING



<110> Brown, Arthur M.  
wible, Barbara A  
Yang, Qing

<120> Protein That Enhances Expression of Potassium Channels on Cell Surfaces and Nucleic Acids That Encode The Same

<130> 22884/04066

<150> 09/062,440

<151> 1998-04-17

<150> 09/712,495

<151> 2000-11-14

<160> 13

<170> PatentIn version 3.1

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<211> 1725

<212> DNA

<213> Rattus norvegicus

<220>

<221> CDS

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| atg | aag | atc | aaa | gaa | ctt | tac | cgc | agg | cgc | ttt | ccc | cgg | aag | acc | ctg | 48 |
| Met | Lys | Ile | Lys | Glu | Leu | Tyr | Arg | Arg | Arg | Phe | Pro | Arg | Lys | Thr | Leu |    |
| 1   |     |     |     | 5   |     |     |     |     | 10  |     |     |     |     | 15  |     |    |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |    |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|----|
| ggg | cct | tcc | gat | ctc | tct | ttg | ctc | tct | ttg | ccc | cct | ggc | acc | tct | cct | 96 |
| Gly | Pro | Ser | Asp | Leu | Ser | Leu | Leu | Ser | Leu | Pro | Pro | Gly | Thr | Ser | Pro |    |
|     |     |     | 20  |     |     |     |     | 25  |     |     |     |     | 30  |     |     |    |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| gta | ggc | tcc | ccc | agc | ccc | ctt | gct | tcc | att | cct | ccc | acc | ctc | ctg | acc | 144 |
| Val | Gly | Ser | Pro | Ser | Pro | Leu | Ala | Ser | Ile | Pro | Pro | Thr | Leu | Leu | Thr |     |
|     |     |     | 35  |     |     |     | 40  |     |     |     |     |     | 45  |     |     |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| cct | ggc | acc | ttg | ctg | ggc | cct | aag | cgt | gag | gtg | gac | atg | cac | cct | cct | 192 |
| Pro | Gly | Thr | Leu | Leu | Gly | Pro | Lys | Arg | Glu | Val | Asp | Met | His | Pro | Pro |     |
|     | 50  |     |     |     |     | 55  |     |     |     |     | 60  |     |     |     |     |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| ctg | ccc | cag | cct | gtg | cac | cct | gac | gtc | acc | atg | aaa | cca | ctg | ccc | ttc | 240 |
| Leu | Pro | Gln | Pro | Val | His | Pro | Asp | Val | Thr | Met | Lys | Pro | Leu | Pro | Phe |     |
| 65  |     |     |     |     | 70  |     |     |     | 75  |     |     |     |     |     | 80  |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| tac | gaa | gtc | tac | gga | gag | ctc | atc | cgg | ccg | acc | acc | ctt | gcg | tcc | acc | 288 |
| Tyr | Glu | Val | Tyr | Gly | Glu | Leu | Ile | Arg | Pro | Thr | Thr | Leu | Ala | Ser | Thr |     |
|     |     |     |     | 85  |     |     |     | 90  |     |     |     |     |     | 95  |     |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| tcc | agt | cag | agg | ttt | gag | gaa | gcc | cac | ttt | acc | ttt | gca | ctc | act | ccc | 336 |
| Ser | Ser | Gln | Arg | Phe | Glu | Glu | Ala | His | Phe | Thr | Phe | Ala | Leu | Thr | Pro |     |
|     |     |     | 100 |     |     |     |     | 105 |     |     |     |     | 110 |     |     |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| cag | cag | ctg | cag | cag | att | ctc | aca | tcc | agg | gag | gtt | ctg | cca | gga | gcc | 384 |
| Gln | Gln | Leu | Gln | Gln | Ile | Leu | Thr | Ser | Arg | Glu | Val | Leu | Pro | Gly | Ala |     |
|     |     | 115 |     |     |     |     | 120 |     |     |     |     | 125 |     |     |     |     |

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|   |      |
|---|------|
| aag tgc gat tat acc ata caa gtg cag ctc agg ttc tgt ctc tgt gag<br>Lys Cys Asp Tyr Thr Ile Gln Val Gln Leu Arg Phe Cys Leu Cys Glu<br>130 135 140     | 432  |
| acc agc tgc ccc cag gag gac tat ttc ccc cct aac ctc ttt gtc aag<br>Thr Ser Cys Pro Gln Glu Asp Tyr Phe Pro Pro Asn Leu Phe Val Lys<br>145 150 155 160 | 480  |
| gtt aat ggg aaa ctc tgc ccc ctg ccg ggt tac ctc cct cca acc aag<br>Val Asn Gly Lys Leu Cys Pro Leu Pro Gly Tyr Leu Pro Pro Thr Lys<br>165 170 175     | 528  |
| aat gga gct gag ccc aag agg cct agt cgt cca atc aac atc aca ccc<br>Asn Gly Ala Glu Pro Lys Arg Pro Ser Arg Pro Ile Asn Ile Thr Pro<br>180 185 190     | 576  |
| ctg gct cgt ctc tca gcc act gtt ccc aac acc ata gtg gtt aac tgg<br>Leu Ala Arg Leu Ser Ala Thr Val Pro Asn Thr Ile Val Val Asn Trp<br>195 200 205     | 624  |
| tca tct gag ttt gga cgg aat tac tcc ttg tct gtg tac ctg gtg agg<br>Ser Ser Glu Phe Gly Arg Asn Tyr Ser Leu Ser Val Tyr Leu Val Arg<br>210 215 220     | 672  |
| cag ttg act gca ggg acc ctg cta caa aag ctc aga gcc aag ggt atc<br>Gln Leu Thr Ala Gly Thr Leu Leu Gln Lys Leu Arg Ala Lys Gly Ile<br>225 230 235 240 | 720  |
| cgg aat cca gac cat tcc cga gca ctg atc aag gag aaa ttg act gct<br>Arg Asn Pro Asp His Ser Arg Ala Leu Ile Lys Glu Lys Leu Thr Ala<br>245 250 255     | 768  |
| gac ccc gac agt gaa gtg gct act aca agt ctc cgg gtg tca ctc atg<br>Asp Pro Asp Ser Glu Val Ala Thr Thr Ser Leu Arg Val Ser Leu Met<br>260 265 270     | 816  |
| tgc ccg ctg ggg aag atg cgc ctg act gtc cca tgc cgc gct ctc acc<br>Cys Pro Leu Gly Lys Met Arg Leu Thr Val Pro Cys Arg Ala Leu Thr<br>275 280 285     | 864  |
| tgt gcc cac ctg cag agt ttc gat gct gcc ctt tat cta cag atg aat<br>Cys Ala His Leu Gln Ser Phe Asp Ala Ala Leu Tyr Leu Gln Met Asn<br>290 295 300     | 912  |
| gag aaa aag cca aca tgg acg tgc cct gtg tgt gac aag aag gct ccc<br>Glu Lys Lys Pro Thr Trp Thr Cys Pro Val Cys Asp Lys Lys Ala Pro<br>305 310 315 320 | 960  |
| tat gag tca ctg att att gat ggt tta ttc atg gaa att ctt aat tcc<br>Tyr Glu Ser Leu Ile Ile Asp Gly Leu Phe Met Glu Ile Leu Asn Ser<br>325 330 335     | 1008 |
| tgt tcg gat tgt gat gag atc cag ttc atg gaa gat gga tcc tgg tgt<br>Cys Ser Asp Cys Asp Glu Ile Gln Phe Met Glu Asp Gly Ser Trp Cys<br>340 345 350     | 1056 |
| cca atg aaa ccc aag aag gag gca tcc gag gtt tgc ccc cca cca ggg<br>Pro Met Lys Pro Lys Lys Glu Ala Ser Glu Val Cys Pro Pro Pro Gly<br>355 360 365     | 1104 |
| tat ggg ctg gat ggt ctc cag tat agc cca gtc cag gag gga aat cag<br>Tyr Gly Leu Asp Gly Leu Gln Tyr Ser Pro Val Gln Glu Gly Asn Gln                    | 1152 |

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380

370

375

|                   |                   |                   |                   |                   |                   |                   |                   |                   |                   |                   |                   |                   |                   |                   |                   |      |
|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|------|
| tca<br>Ser<br>385 | gag<br>Glu        | aat<br>Asn        | aag<br>Lys        | aag<br>Lys        | agg<br>Arg<br>390 | gtt<br>Val        | gaa<br>Glu        | gtc<br>Val        | att<br>Ile        | gac<br>Asp<br>395 | ttg<br>Leu        | aca<br>Thr        | atc<br>Ile        | gaa<br>Glu        | agc<br>Ser<br>400 | 1200 |
| tca<br>Ser        | tca<br>Ser        | gat<br>Asp        | gag<br>Glu        | gaa<br>Glu<br>405 | gat<br>Asp        | ctg<br>Leu        | ccc<br>Pro        | ccc<br>Pro        | acc<br>Thr<br>410 | aag<br>Lys        | aag<br>Lys        | cac<br>His        | tgc<br>Cys        | cct<br>Pro<br>415 | gtt<br>Val        | 1248 |
| acc<br>Thr        | tcg<br>Ser        | gct<br>Ala        | gcc<br>Ala<br>420 | att<br>Ile        | cca<br>Pro        | gcc<br>Ala        | ctt<br>Leu        | cct<br>Pro<br>425 | gga<br>Gly        | agc<br>Ser        | aaa<br>Lys        | gga<br>Gly        | gcc<br>Ala<br>430 | ctg<br>Leu        | acc<br>Thr        | 1296 |
| tct<br>Ser        | ggt<br>Gly        | cac<br>His<br>435 | cag<br>Gln        | ccg<br>Pro        | tct<br>Ser        | tcg<br>Ser        | gtg<br>Val<br>440 | ctg<br>Leu        | cgg<br>Arg        | agc<br>Ser        | cct<br>Pro        | gca<br>Ala<br>445 | atg<br>Met        | ggt<br>Gly        | aca<br>Thr        | 1344 |
| ctg<br>Leu        | ggc<br>Gly<br>450 | agt<br>Ser        | gat<br>Asp        | ttc<br>Phe        | ctg<br>Leu        | tct<br>Ser<br>455 | agt<br>Ser        | ctc<br>Leu        | cca<br>Pro        | cta<br>Leu        | cat<br>His<br>460 | gag<br>Glu        | tac<br>Tyr        | cca<br>Pro        | cct<br>Pro        | 1392 |
| gcc<br>Ala<br>465 | ttc<br>Phe        | ccg<br>Pro        | ctg<br>Leu        | ggg<br>Gly        | gct<br>Ala<br>470 | gac<br>Asp        | atc<br>Ile        | caa<br>Gln        | ggt<br>Gly        | tta<br>Leu<br>475 | gat<br>Asp        | tta<br>Leu        | ttt<br>Phe        | tct<br>Ser        | ttc<br>Phe<br>480 | 1440 |
| ctt<br>Leu        | cag<br>Gln        | act<br>Thr        | gag<br>Glu        | agt<br>Ser<br>485 | cag<br>Gln        | cac<br>His        | tac<br>Tyr        | agc<br>Ser        | cct<br>Pro<br>490 | tca<br>Ser        | gtt<br>Val        | atc<br>Ile        | act<br>Thr        | tca<br>Ser<br>495 | cta<br>Leu        | 1488 |
| gat<br>Asp        | gag<br>Glu        | cag<br>Gln        | gac<br>Asp<br>500 | acc<br>Thr        | ctt<br>Leu        | ggc<br>Gly        | cac<br>His        | ttc<br>Phe<br>505 | ttc<br>Phe        | caa<br>Gln        | ttc<br>Phe        | cgg<br>Arg        | gga<br>Gly<br>510 | acc<br>Thr        | cct<br>Pro        | 1536 |
| ccc<br>Pro        | cac<br>His        | ttc<br>Phe<br>515 | ctg<br>Leu        | ggc<br>Gly        | cca<br>Pro        | ctg<br>Leu        | gcc<br>Ala<br>520 | ccc<br>Pro        | aca<br>Thr        | ttg<br>Leu        | ggg<br>Gly        | agc<br>Ser<br>525 | tct<br>Ser        | cac<br>His        | cgc<br>Arg        | 1584 |
| agc<br>Ser        | gcc<br>Ala<br>530 | act<br>Thr        | cca<br>Pro        | gca<br>Ala        | ccc<br>Pro        | gct<br>Ala<br>535 | cct<br>Pro        | ggc<br>Gly        | cgt<br>Arg        | gtc<br>Val        | agc<br>Ser<br>540 | agc<br>Ser        | att<br>Ile        | gtg<br>Val        | gct<br>Ala        | 1632 |
| cct<br>Pro<br>545 | ggg<br>Gly        | agt<br>Ser        | tcc<br>Ser        | ttg<br>Leu        | agg<br>Arg<br>550 | gaa<br>Glu        | ggg<br>Gly        | cat<br>His        | gga<br>Gly        | gga<br>Gly<br>555 | ccc<br>Pro        | ctg<br>Leu        | cct<br>Pro        | tcc<br>Ser        | ggt<br>Gly<br>560 | 1680 |
| ccc<br>Pro        | tct<br>Ser        | ttg<br>Leu        | act<br>Thr        | ggc<br>Gly<br>565 | tgt<br>Cys        | cgg<br>Arg        | tca<br>Ser        | gac<br>Asp        | gtc<br>Val<br>570 | att<br>Ile        | tcc<br>Ser        | ttg<br>Leu        | gac<br>Asp        | tga               |                   | 1725 |

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Gly Pro Ser Asp Leu Ser Leu Leu Ser Leu Pro Pro Gly Thr Ser Pro  
Page 3

Val Gly Ser Pro Ser Pro Leu Ala Ser Ile Pro Pro Thr Leu Leu Thr  
 35 40 45  
 Pro Gly Thr Leu Leu Gly Pro Lys Arg Glu Val Asp Met His Pro Pro  
 50 55 60  
 Leu Pro Gln Pro Val His Pro Asp Val Thr Met Lys Pro Leu Pro Phe  
 65 70 75 80  
 Tyr Glu Val Tyr Gly Glu Leu Ile Arg Pro Thr Thr Leu Ala Ser Thr  
 85 90 95  
 Ser Ser Gln Arg Phe Glu Glu Ala His Phe Thr Phe Ala Leu Thr Pro  
 100 105 110  
 Gln Gln Leu Gln Gln Ile Leu Thr Ser Arg Glu Val Leu Pro Gly Ala  
 115 120 125  
 Lys Cys Asp Tyr Thr Ile Gln Val Gln Leu Arg Phe Cys Leu Cys Glu  
 130 135 140  
 Thr Ser Cys Pro Gln Glu Asp Tyr Phe Pro Pro Asn Leu Phe Val Lys  
 145 150 155 160  
 Val Asn Gly Lys Leu Cys Pro Leu Pro Gly Tyr Leu Pro Pro Thr Lys  
 165 170 175  
 Asn Gly Ala Glu Pro Lys Arg Pro Ser Arg Pro Ile Asn Ile Thr Pro  
 180 185 190  
 Leu Ala Arg Leu Ser Ala Thr Val Pro Asn Thr Ile Val Val Asn Trp  
 195 200 205  
 Ser Ser Glu Phe Gly Arg Asn Tyr Ser Leu Ser Val Tyr Leu Val Arg  
 210 215 220  
 Gln Leu Thr Ala Gly Thr Leu Leu Gln Lys Leu Arg Ala Lys Gly Ile  
 225 230 235 240  
 Arg Asn Pro Asp His Ser Arg Ala Leu Ile Lys Glu Lys Leu Thr Ala  
 245 250 255  
 Asp Pro Asp Ser Glu Val Ala Thr Thr Ser Leu Arg Val Ser Leu Met  
 260 265 270

Cys Pro Leu Gly Lys Met Arg Leu Thr Val Pro Cys Arg Ala Leu Thr  
 275 280 285  
 Cys Ala His Leu Gln Ser Phe Asp Ala Ala Leu Tyr Leu Gln Met Asn  
 290 295 300  
 Glu Lys Lys Pro Thr Trp Thr Cys Pro Val Cys Asp Lys Lys Ala Pro  
 305 310 315 320  
 Tyr Glu Ser Leu Ile Ile Asp Gly Leu Phe Met Glu Ile Leu Asn Ser  
 325 330 335  
 Cys Ser Asp Cys Asp Glu Ile Gln Phe Met Glu Asp Gly Ser Trp Cys  
 340 345 350  
 Pro Met Lys Pro Lys Lys Glu Ala Ser Glu Val Cys Pro Pro Pro Gly  
 355 360 365  
 Tyr Gly Leu Asp Gly Leu Gln Tyr Ser Pro Val Gln Glu Gly Asn Gln  
 370 375 380  
 Ser Glu Asn Lys Lys Arg Val Glu Val Ile Asp Leu Thr Ile Glu Ser  
 385 390 395 400  
 Ser Ser Asp Glu Glu Asp Leu Pro Pro Thr Lys Lys His Cys Pro Val  
 405 410 415  
 Thr Ser Ala Ala Ile Pro Ala Leu Pro Gly Ser Lys Gly Ala Leu Thr  
 420 425 430  
 Ser Gly His Gln Pro Ser Ser Val Leu Arg Ser Pro Ala Met Gly Thr  
 435 440 445  
 Leu Gly Ser Asp Phe Leu Ser Ser Leu Pro Leu His Glu Tyr Pro Pro  
 450 455 460  
 Ala Phe Pro Leu Gly Ala Asp Ile Gln Gly Leu Asp Leu Phe Ser Phe  
 465 470 475 480  
 Leu Gln Thr Glu Ser Gln His Tyr Ser Pro Ser Val Ile Thr Ser Leu  
 485 490 495  
 Asp Glu Gln Asp Thr Leu Gly His Phe Phe Gln Phe Arg Gly Thr Pro  
 500 505 510  
 Pro His Phe Leu Gly Pro Leu Ala Pro Thr Leu Gly Ser Ser His Arg  
 515 520 525

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Ser Ala Thr Pro Ala Pro Ala Pro Gly Arg Val Ser Ser Ile Val Ala  
530 535 540

Pro Gly Ser Ser Leu Arg Glu Gly His Gly Gly Pro Leu Pro Ser Gly  
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Pro Ser Leu Thr Gly Cys Arg Ser Asp Val Ile Ser Leu Asp  
565 570

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1 5 10 15

ggg ccc tct gat ctc tcc ctt ctc tct ttg ccc cct ggc acc tct cct 96  
Gly Pro Ser Asp Leu Ser Leu Leu Ser Leu Pro Pro Gly Thr Ser Pro  
20 25 30

gta ggc tcc cct ggt cct cta gct ccc att ccc cca acg ctg ttg gcc 144  
Val Gly Ser Pro Gly Pro Leu Ala Pro Ile Pro Pro Thr Leu Leu Ala  
35 40 45

cct ggc acc ctg ctg ggc ccc aag cgt gag gtg gac atg cac ccc cct 192  
Pro Gly Thr Leu Leu Gly Pro Lys Arg Glu Val Asp Met His Pro Pro  
50 55 60

ctg ccc cag cct gtg cac cct gat gtc acc atg aaa cca ttg ccc ttc 240  
Leu Pro Gln Pro Val His Pro Asp Val Thr Met Lys Pro Leu Pro Phe  
65 70 75 80

tat gaa gtc tat ggg gag ctc atc cgg ccc acc acc ctt gca tcc act 288  
Tyr Glu Val Tyr Gly Glu Leu Ile Arg Pro Thr Thr Leu Ala Ser Thr  
85 90 95

tct agc cag cgg ttt gag gaa gcg cac ttt acc ttt gcc ctc aca ccc 336  
Ser Ser Gln Arg Phe Glu Glu Ala His Phe Thr Phe Ala Leu Thr Pro  
100 105 110

cag caa gtg cag cag att ctt aca tcc aga gag gtt ctg cca gga gcc 384  
Gln Gln Val Gln Gln Ile Leu Thr Ser Arg Glu Val Leu Pro Gly Ala  
115 120 125

aaa tgt gat tat acc ata cag gtg cag cta agg ttc tgt ctc tgt gag 432  
Lys Cys Asp Tyr Thr Ile Gln Val Gln Leu Arg Phe Cys Leu Cys Glu  
130 135 140

acc agc tgc ccc cag gaa gat tat ttt ccc ccc aac ctc ttt gtc aag 480  
Thr Ser Cys Pro Gln Glu Asp Tyr Phe Pro Pro Asn Leu Phe Val Lys  
145 150 155 160

## 228844066.ST25.txt

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| gtt aat ggg aaa ctg tgc ccc ctg ccg ggt tac ctt ccc cca acc aag<br>Val Asn Gly Lys Leu Cys Pro Leu Pro Gly Tyr Leu Pro Pro Thr Lys<br>165 170 175     | 528  |
| aat ggg gcc gag ccc aag agg ccc agc cgc ccc atc aac atc aca ccc<br>Asn Gly Ala Glu Pro Lys Arg Pro Ser Arg Pro Ile Asn Ile Thr Pro<br>180 185 190     | 576  |
| ctg gct cga ctc tca gcc act gtt ccc aac acc att gtg gtc aat tgg<br>Leu Ala Arg Leu Ser Ala Thr Val Pro Asn Thr Ile Val Val Asn Trp<br>195 200 205     | 624  |
| tca tct gag ttc gga cgg aat tac tcc ttg tct gtg tac ctg gtg agg<br>Ser Ser Glu Phe Gly Arg Asn Tyr Ser Leu Ser Val Tyr Leu Val Arg<br>210 215 220     | 672  |
| cag ttg act gca gga acc ctt cta caa aaa ctc aga gca aag ggt atc<br>Gln Leu Thr Ala Gly Thr Leu Leu Gln Lys Leu Arg Ala Lys Gly Ile<br>225 230 235 240 | 720  |
| cgg aac cca gac cac tcg cgg gca ctg atc aag gag aaa ttg act gct<br>Arg Asn Pro Asp His Ser Arg Ala Leu Ile Lys Glu Lys Leu Thr Ala<br>245 250 255     | 768  |
| gac cct gac agt gag gtg gcc act aca agt ctc cgg gtg tca ctc atg<br>Asp Pro Asp Ser Glu Val Ala Thr Thr Ser Leu Arg Val Ser Leu Met<br>260 265 270     | 816  |
| tgc ccg cta ggg aag atg cgc ctg act gtc cct tgt cgt gcc ctc acc<br>Cys Pro Leu Gly Lys Met Arg Leu Thr Val Pro Cys Arg Ala Leu Thr<br>275 280 285     | 864  |
| tgt gcc cac ctg cag agc ttc gat gct gcc ctt tat cta cag atg aat<br>Cys Ala His Leu Gln Ser Phe Asp Ala Ala Leu Tyr Leu Gln Met Asn<br>290 295 300     | 912  |
| gag aag aag cct aca tgg aca tgt cct gtg tgt gac aag aag gct ccc<br>Glu Lys Lys Pro Thr Trp Thr Cys Pro Val Cys Asp Lys Lys Ala Pro<br>305 310 315 320 | 960  |
| tat gaa tct ctt atc att gat ggt tta ttt atg gag att ctt agt tcc<br>Tyr Glu Ser Leu Ile Ile Asp Gly Leu Phe Met Glu Ile Leu Ser Ser<br>325 330 335     | 1008 |
| tgt tca gat tgt gat gag atc caa ttc atg gaa gat gga tcc tgg tgc<br>Cys Ser Asp Cys Asp Glu Ile Gln Phe Met Glu Asp Gly Ser Trp Cys<br>340 345 350     | 1056 |
| cca atg aaa ccc aag aag gag gca tct gag gtt tgc ccc ccg cca ggg<br>Pro Met Lys Pro Lys Lys Glu Ala Ser Glu Val Cys Pro Pro Gly<br>355 360 365         | 1104 |
| tat ggg ctg gat ggc ctc cag tac agc cca gtc cag ggg gga gat cca<br>Tyr Gly Leu Asp Gly Leu Gln Tyr Ser Pro Val Gln Gly Gly Asp Pro<br>370 375 380     | 1152 |
| tca gag aat aag aag aag gtc gaa gtt att gac ttg aca ata gaa agc<br>Ser Glu Asn Lys Lys Lys Val Glu Val Ile Asp Leu Thr Ile Glu Ser<br>385 390 395 400 | 1200 |
| tca tca gat gag gag gat ctg ccc cct acc aag aag cac tgt tct gtc<br>Ser Ser Asp Glu Glu Asp Leu Pro Pro Thr Lys Lys His Cys Ser Val<br>405 410 415     | 1248 |

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405

410

415

|   |      |
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| Thr Ser Ala Ala Ile Pro Ala Leu Pro Gly Ser Lys Gly Val Leu Thr |      |
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| tct ggc cac cag cca tcc tcg gtg cta agg agc cct gct atg ggc acg | 1344 |
| Ser Gly His Gln Pro Ser Ser Val Leu Arg Ser Pro Ala Met Gly Thr |      |
| 435 440 445   |      |
| ttg ggt ggg gat ttc ctg tcc agt ctc cca cta cat gag tac cca cct | 1392 |
| Leu Gly Gly Asp Phe Leu Ser Ser Leu Pro Leu His Glu Tyr Pro Pro |      |
| 450 455 460   |      |
| gcc ttc cca ctg gga gcc gac atc caa ggt tta gat tta ttt tca ttt | 1440 |
| Ala Phe Pro Leu Gly Ala Asp Ile Gln Gly Leu Asp Leu Phe Ser Phe |      |
| 465 470 475 480   |      |
| ctt cag aca gag agt cag cac tat ggc ccc tct gtc atc acc tca cta | 1488 |
| Leu Gln Thr Glu Ser Gln His Tyr Gly Pro Ser Val Ile Thr Ser Leu |      |
| 485 490 495   |      |
| gat gaa cag gat gcc ctt ggc cac ttc ttc cag tac cga ggg acc cct | 1536 |
| Asp Glu Gln Asp Ala Leu Gly His Phe Phe Gln Tyr Arg Gly Thr Pro |      |
| 500 505 510   |      |
| tct cac ttt ctg ggc cca ctg gcc ccc acg ctg ggg agc tcc cac tgc | 1584 |
| Ser His Phe Leu Gly Pro Leu Ala Pro Thr Leu Gly Ser Ser His Cys |      |
| 515 520 525   |      |
| agc gcc act ccg gcg ccc cct cct ggc cgt gtc agc agc att gtg gcc | 1632 |
| Ser Ala Thr Pro Ala Pro Pro Gly Arg Val Ser Ser Ile Val Ala     |      |
| 530 535 540   |      |
| cct ggg ggg gcc ttg agg gag ggg cat gga gga ccc ctg ccc tca ggt | 1680 |
| Pro Gly Gly Ala Leu Arg Glu Gly His Gly Gly Pro Leu Pro Ser Gly |      |
| 545 550 555 560   |      |
| ccc tct ttg act ggc tgt cgg tca gac atc att tcc ctg gac tga     | 1725 |
| Pro Ser Leu Thr Gly Cys Arg Ser Asp Ile Ile Ser Leu Asp         |      |
| 565 570   |      |

<210> 4  
 <211> 574  
 <212> PRT  
 <213> Homo sapiens

<400> 4

Met Lys Ile Lys Glu Leu Tyr Arg Arg Arg Phe Pro Arg Lys Thr Leu  
 1 5 10 15

Gly Pro Ser Asp Leu Ser Leu Leu Ser Leu Pro Pro Gly Thr Ser Pro  
 20 25 30

Val Gly Ser Pro Gly Pro Leu Ala Pro Ile Pro Pro Thr Leu Leu Ala  
 35 40 45

Pro Gly Thr Leu Leu Gly Pro Lys Arg Glu Val Asp Met His Pro Pro  
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50

55

60

Leu Pro Gln Pro Val His Pro Asp Val Thr Met Lys Pro Leu Pro Phe  
65 70 75 80

Tyr Glu Val Tyr Gly Glu Leu Ile Arg Pro Thr Thr Leu Ala Ser Thr  
85 90 95

Ser Ser Gln Arg Phe Glu Glu Ala His Phe Thr Phe Ala Leu Thr Pro  
100 105 110

Gln Gln Val Gln Gln Ile Leu Thr Ser Arg Glu Val Leu Pro Gly Ala  
115 120 125

Lys Cys Asp Tyr Thr Ile Gln Val Gln Leu Arg Phe Cys Leu Cys Glu  
130 135 140

Thr Ser Cys Pro Gln Glu Asp Tyr Phe Pro Pro Asn Leu Phe Val Lys  
145 150 155 160

Val Asn Gly Lys Leu Cys Pro Leu Pro Gly Tyr Leu Pro Pro Thr Lys  
165 170 175

Asn Gly Ala Glu Pro Lys Arg Pro Ser Arg Pro Ile Asn Ile Thr Pro  
180 185 190

Leu Ala Arg Leu Ser Ala Thr Val Pro Asn Thr Ile Val Val Asn Trp  
195 200 205

Ser Ser Glu Phe Gly Arg Asn Tyr Ser Leu Ser Val Tyr Leu Val Arg  
210 215 220

Gln Leu Thr Ala Gly Thr Leu Leu Gln Lys Leu Arg Ala Lys Gly Ile  
225 230 235 240

Arg Asn Pro Asp His Ser Arg Ala Leu Ile Lys Glu Lys Leu Thr Ala  
245 250 255

Asp Pro Asp Ser Glu Val Ala Thr Thr Ser Leu Arg Val Ser Leu Met  
260 265 270

Cys Pro Leu Gly Lys Met Arg Leu Thr Val Pro Cys Arg Ala Leu Thr  
275 280 285

Cys Ala His Leu Gln Ser Phe Asp Ala Ala Leu Tyr Leu Gln Met Asn  
290 295 300

## 228844066.ST25.txt

Glu Lys Lys Pro Thr Trp Thr Cys Pro Val Cys Asp Lys Lys Ala Pro  
 305 310 315 320

Tyr Glu Ser Leu Ile Ile Asp Gly Leu Phe Met Glu Ile Leu Ser Ser  
 325 330 335

Cys Ser Asp Cys Asp Glu Ile Gln Phe Met Glu Asp Gly Ser Trp Cys  
 340 345 350

Pro Met Lys Pro Lys Lys Glu Ala Ser Glu Val Cys Pro Pro Pro Gly  
 355 360 365

Tyr Gly Leu Asp Gly Leu Gln Tyr Ser Pro Val Gln Gly Gly Asp Pro  
 370 375 380

Ser Glu Asn Lys Lys Lys Val Glu Val Ile Asp Leu Thr Ile Glu Ser  
 385 390 395 400

Ser Ser Asp Glu Glu Asp Leu Pro Pro Thr Lys Lys His Cys Ser Val  
 405 410 415

Thr Ser Ala Ala Ile Pro Ala Leu Pro Gly Ser Lys Gly Val Leu Thr  
 420 425 430

Ser Gly His Gln Pro Ser Ser Val Leu Arg Ser Pro Ala Met Gly Thr  
 435 440 445

Leu Gly Gly Asp Phe Leu Ser Ser Leu Pro Leu His Glu Tyr Pro Pro  
 450 455 460

Ala Phe Pro Leu Gly Ala Asp Ile Gln Gly Leu Asp Leu Phe Ser Phe  
 465 470 475 480

Leu Gln Thr Glu Ser Gln His Tyr Gly Pro Ser Val Ile Thr Ser Leu  
 485 490 495

Asp Glu Gln Asp Ala Leu Gly His Phe Phe Gln Tyr Arg Gly Thr Pro  
 500 505 510

Ser His Phe Leu Gly Pro Leu Ala Pro Thr Leu Gly Ser Ser His Cys  
 515 520 525

Ser Ala Thr Pro Ala Pro Pro Pro Gly Arg Val Ser Ser Ile Val Ala  
 530 535 540

Pro Gly Gly Ala Leu Arg Glu Gly His Gly Gly Pro Leu Pro Ser Gly  
 545 550 555 560

228844066.ST25.txt

Pro Ser Leu Thr Gly Cys Arg Ser Asp Ile Ile Ser Leu Asp  
565 570

<210> 5  
<211> 99  
<212> PRT  
<213> Homo sapiens

<400> 5

Thr Trp Thr Cys Pro Val Cys Asp Lys Lys Ala Pro Tyr Glu Ser Leu  
1 5 10 15

Ile Ile Asp Gly Leu Phe Met Glu Ile Leu Ser Ser Cys Ser Asp Cys  
20 25 30

Asp Glu Ile Gln Phe Met Glu Asp Gly Ser Trp Cys Pro Met Lys Pro  
35 40 45

Lys Lys Glu Ala Ser Glu Val Cys Pro Pro Pro Gly Tyr Gly Leu Asp  
50 55 60

Gly Leu Gln Tyr Ser Pro Val Gln Gly Gly Asp Pro Ser Glu Asn Lys  
65 70 75 80

Lys Lys Val Glu Val Ile Asp Leu Thr Ile Glu Ser Ser Ser Asp Glu  
85 90 95

Glu Asp Leu

<210> 6  
<211> 574  
<212> PRT  
<213> Synthetic construct

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<222> (37)..(37)  
<223> xaa = glycine or serine

<220>  
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<223> xaa = proline or serine

<220>  
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<222> (48)..(48)  
<223> xaa = alanine or threonine

<220>  
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<222> (115)..(115)  
<223> Xaa = valine or leucine

<220>  
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<222> (335)..(335)  
<223> Xaa = serine or asparagine

<220>  
<221> MISC\_FEATURE  
<222> (381)..(381)  
<223> Xaa = glycine or glutamic acid

<220>  
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<222> (383)..(383)  
<223> Xaa = aspartic acid or asparagine

<220>  
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<223> Xaa = isoproline or glutamine

<220>  
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<222> (390)..(390)  
<223> Xaa = lysine or arginine

<220>  
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<223> Xaa = serine or proline

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<223> Xaa = valine or alanine

<220>  
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<222> (451)..(451)  
<223> Xaa = glycine or serine

<220>  
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<222> (489)..(489)  
<223> Xaa = glycine or serine

<220>  
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<223> Xaa = alanine or threonine

<220>

<221> MISC\_FEATURE

<222> (508)..(508)

<223> Xaa = tyrosine or phenylalanine

<220>

<221> MISC\_FEATURE

<222> (513)..(513)

<223> Xaa = serine or proline

<220>

<221> MISC\_FEATURE

<222> (528)..(528)

<223> Xaa = cysteine or arginine

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<222> (535)..(535)

<223> Xaa = proline or alanine

<220>

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<222> (547)..(547)

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<220>

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<222> (571)..(571)

<223> Xaa = isoleucine or valine

<400> 6

Met Lys Ile Lys Glu Leu Tyr Arg Arg Arg Phe Pro Arg Lys Thr Leu  
1 5 10 15

Gly Pro Ser Asp Leu Ser Leu Leu Ser Leu Pro Pro Gly Thr Ser Pro  
20 25 30

Val Gly Ser Pro Xaa Pro Leu Ala Xaa Ile Pro Pro Thr Leu Leu Xaa  
35 40 45

Pro Gly Thr Leu Leu Gly Pro Lys Arg Glu Val Asp Met His Pro Pro  
50 55 60

228844066.ST25.txt

Leu Pro Gln Pro Val His Pro Asp Val Thr Met Lys Pro Leu Pro Phe  
 65 70 75 80  
 Tyr Glu Val Tyr Gly Glu Leu Ile Arg Pro Thr Thr Leu Ala Ser Thr  
 85 90 95  
 Ser Ser Gln Arg Phe Glu Glu Ala His Phe Thr Phe Ala Leu Thr Pro  
 100 105 110  
 Gln Gln Xaa Gln Gln Ile Leu Thr Ser Arg Glu Val Leu Pro Gly Ala  
 115 120 125  
 Lys Cys Asp Tyr Thr Ile Gln Val Gln Leu Arg Phe Cys Leu Cys Glu  
 130 135 140  
 Thr Ser Cys Pro Gln Glu Asp Tyr Phe Pro Pro Asn Leu Phe Val Lys  
 145 150 155 160  
 Val Asn Gly Lys Leu Cys Pro Leu Pro Gly Tyr Leu Pro Pro Thr Lys  
 165 170 175  
 Asn Gly Ala Glu Pro Lys Arg Pro Ser Arg Pro Ile Asn Ile Thr Pro  
 180 185 190  
 Leu Ala Arg Leu Ser Ala Thr Val Pro Asn Thr Ile Val Val Asn Trp  
 195 200 205  
 Ser Ser Glu Phe Gly Arg Asn Tyr Ser Leu Ser Val Tyr Leu Val Arg  
 210 215 220  
 Gln Leu Thr Ala Gly Thr Leu Leu Gln Lys Leu Arg Ala Lys Gly Ile  
 225 230 235 240  
 Arg Asn Pro Asp His Ser Arg Ala Leu Ile Lys Glu Lys Leu Thr Ala  
 245 250 255  
 Asp Pro Asp Ser Glu Val Ala Thr Thr Ser Leu Arg Val Ser Leu Met  
 260 265 270  
 Cys Pro Leu Gly Lys Met Arg Leu Thr Val Pro Cys Arg Ala Leu Thr  
 275 280 285  
 Cys Ala His Leu Gln Ser Phe Asp Ala Ala Leu Tyr Leu Gln Met Asn  
 290 295 300  
 Glu Lys Lys Pro Thr Trp Thr Cys Pro Val Cys Asp Lys Lys Ala Pro  
 305 310 315 320

228844066.ST25.txt

Tyr Glu Ser Leu Ile Ile Asp Gly Leu Phe Met Glu Ile Leu Xaa Ser  
 325 330 335  
 Cys Ser Asp Cys Asp Glu Ile Gln Phe Met Glu Asp Gly Ser Trp Cys  
 340 345 350  
 Pro Met Lys Pro Lys Lys Glu Ala Ser Glu Val Cys Pro Pro Pro Gly  
 355 360 365  
 Tyr Gly Leu Asp Gly Leu Gln Tyr Ser Pro Val Gln Xaa Gly Xaa Pro  
 370 375 380  
 Ser Glu Asn Lys Lys Xaa Val Glu Val Ile Asp Leu Thr Ile Glu Ser  
 385 390 395 400  
 Ser Ser Asp Glu Glu Asp Leu Pro Pro Thr Lys Lys His Cys Xaa Val  
 405 410 415  
 Thr Ser Ala Ala Ile Pro Ala Leu Pro Gly Ser Lys Gly Xaa Leu Thr  
 420 425 430  
 Ser Gly His Gln Pro Ser Ser Val Leu Arg Ser Pro Ala Met Gly Thr  
 435 440 445  
 Leu Gly Xaa Asp Phe Leu Ser Ser Leu Pro Leu His Glu Tyr Pro Pro  
 450 455 460  
 Ala Phe Pro Leu Gly Ala Asp Ile Gln Gly Leu Asp Leu Phe Ser Phe  
 465 470 475 480  
 Leu Gln Thr Glu Ser Gln His Tyr Xaa Pro Ser Val Ile Thr Ser Leu  
 485 490 495  
 Asp Glu Gln Asp Xaa Leu Gly His Phe Phe Gln Xaa Arg Gly Thr Pro  
 500 505 510  
 Xaa His Phe Leu Gly Pro Leu Ala Pro Thr Leu Gly Ser Ser His Xaa  
 515 520 525  
 Ser Ala Thr Pro Ala Pro Xaa Pro Gly Arg Val Ser Ser Ile Val Ala  
 530 535 540  
 Pro Gly Xaa Xaa Leu Arg Glu Gly His Gly Gly Pro Leu Pro Ser Gly  
 545 550 555 560  
 Pro Ser Leu Thr Gly Cys Arg Ser Asp Ile Xaa Ser Leu Asp  
 565 570

<210> 7  
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 <213> Rattus norvegicus

<400> 7

Thr Trp Thr Cys Pro Val Cys Asp Lys Lys Ala Pro Tyr Glu Ser Leu  
 1 5 10 15

Ile Ile Asp Gly Leu Phe Met Glu Ile Leu Asn Ser Cys Ser Asp Cys  
 20 25 30

Asp Glu Ile Gln Phe Met Glu Asp Gly Ser Trp Cys Pro Met Lys Pro  
 35 40 45

Lys Lys Glu Ala Ser Glu Val Cys Pro Pro Pro Gly Tyr Gly Leu Asp  
 50 55 60

Gly Leu Gln Tyr Ser Pro Val Gln Glu Gly Asn Gln Ser Glu Asn Lys  
 65 70 75 80

Lys Arg Val Glu Val Ile Asp Leu Thr Ile Glu Ser Ser Ser Asp Glu  
 85 90 95

Glu Asp Leu

<210> 8  
 <211> 167  
 <212> PRT  
 <213> Homo sapiens

<400> 8

Pro Pro Thr Lys Lys His Cys Ser Val Thr Ser Ala Ala Ile Pro Ala  
 1 5 10 15

Leu Pro Gly Ser Lys Gly Val Leu Thr Ser Gly His Gln Pro Ser Ser  
 20 25 30

Val Leu Arg Ser Pro Ala Met Gly Thr Leu Gly Gly Asp Phe Leu Ser  
 35 40 45

Ser Leu Pro Leu His Glu Tyr Pro Pro Ala Phe Pro Leu Gly Ala Asp  
 50 55 60

Ile Gln Gly Leu Asp Leu Phe Ser Phe Leu Gln Thr Glu Ser Gln His  
 65 70 75 80



228844066.ST25.txt

Tyr Gly Pro Ser Val Ile Thr Ser Leu Asp Glu Gln Asp Ala Leu Gly  
85 90 95

His Phe Phe Gln Tyr Arg Gly Thr Pro Ser His Phe Leu Gly Pro Leu  
100 105 110

Ala Pro Thr Leu Gly Ser Ser His Cys Ser Ala Thr Pro Ala Pro Pro  
115 120 125

Pro Gly Ala Val Ser Ser Ile Val Ala Pro Gly Gly Ala Leu Arg Glu  
130 135 140

Gly His Gly Gly Pro Leu Pro Ser Gly Pro Ser Leu Thr Gly Cys Arg  
145 150 155 160

Ser Asp Ile Ile Ser Leu Asp  
165

<210> 9  
<211> 167  
<212> PRT  
<213> Rattus norvegicus

<400> 9

Pro Pro Thr Lys Lys His Cys Pro Val Thr Ser Ala Ala Ile Pro Ala  
1 5 10 15

Leu Pro Gly Ser Lys Gly Ala Leu Thr Ser Gly His Gln Pro Ser Ser  
20 25 30

Val Leu Arg Ser Pro Ala Met Gly Thr Leu Gly Ser Asp Phe Leu Ser  
35 40 45

Ser Leu Pro Leu His Glu Tyr Pro Pro Ala Phe Pro Leu Gly Ala Asp  
50 55 60

Ile Gln Gly Leu Asp Leu Phe Ser Phe Leu Gln Thr Glu Ser Gln His  
65 70 75 80

Tyr Ser Pro Ser Val Ile Thr Ser Leu Asp Glu Gln Asp Thr Leu Gly  
85 90 95

His Phe Phe Gln Phe Arg Gly Thr Pro Pro His Phe Leu Gly Pro Leu  
100 105 110

Ala Pro Thr Leu Gly Ser Ser His Arg Ser Ala Thr Pro Ala Pro Ala  
115 120 125

228844066.ST25.txt

Pro Gly Arg Val Ser Ser Ile Val Ala Pro Gly Ser Ser Leu Arg Glu  
130 135 140

Gly His Gly Gly Pro Leu Pro Ser Gly Pro Ser Leu Thr Gly Cys Arg  
145 150 155 160

Ser Asp Val Ile Ser Leu Asp  
165

<210> 10  
<211> 98  
<212> PRT  
<213> synthetic construct

<220>  
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<223> Xaa = serine or asparagine

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<222> (72)..(72)  
<223> Xaa = glycine or glutamic acid

<220>  
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<222> (74)..(74)  
<223> Xaa = aspartic acid or asparagine

<220>  
<221> misc\_feature  
<222> (75)..(75)  
<223> Xaa = proline or glutamine

<220>  
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<222> (81)..(81)  
<223> Xaa = lysine or arginine

<400> 10

Trp Thr Cys Pro Val Cys Asp Lys Lys Ala Pro Tyr Glu Ser Leu Ile  
1 5 10 15

Ile Asp Gly Leu Phe Met Glu Ile Leu Xaa Ser Cys Ser Asp Cys Asp  
20 25 30

Glu Ile Gln Phe Met Glu Asp Gly Ser Trp Cys Pro Met Lys Pro Lys  
35 40 45

Lys Glu Ala Ser Glu Val Cys Pro Pro Pro Gly Tyr Gly Leu Asp Gly

50

55

Leu Gln Tyr Ser Pro Val Gln Xaa Gly Xaa Xaa Ser Glu Asn Lys Lys  
65 70 75 80

xaa val Glu val Ile Asp Leu Thr Ile Glu Ser Ser Ser Asp Glu Glu  
85 90 95

Asp Leu

<210> 11  
<211> 167  
<212> PRT  
<213> Synthetic construct

<220>  
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<223> Xaa = serine or proline

<220>  
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<222> (23)..(23)  
<223> Xaa = valine or alanine

<220>  
<221> misc\_feature  
<222> (44)..(44)  
<223> Xaa = glycine or serine

<220>  
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<222> (82)..(82)  
<223> Xaa = glycine or serine

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<222> (94)..(94)  
<223> Xaa = alanine or threonine

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<222> (101)..(101)  
<223> Xaa = tyrosine or phenylalanine

<220>  
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<222> (106)..(106)  
<223> Xaa = serine or proline

<220>

<221> misc\_feature  
 <222> (121)..(121)  
 <223> Xaa = cysteine or alanine

<220>  
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 <222> (128)..(128)  
 <223> Xaa = proline or alanine

<220>  
 <221> misc\_feature  
 <222> (140)..(140)  
 <223> Xaa = glycine or serine

<220>  
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 <222> (141)..(141)  
 <223> Xaa = alanine or serine

<220>  
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 <222> (164)..(164)  
 <223> Xaa = isoleucine or valine

<400> 11

Pro Pro Thr Lys Lys His Cys Xaa Val Thr Ser Ala Ala Ile Pro Ala  
 1 5 10 15

Leu Pro Gly Ser Lys Gly Xaa Leu Thr Ser Gly His Gln Pro Ser Ser  
 20 25 30

Val Leu Arg Ser Pro Ala Met Gly Thr Leu Gly Xaa Asp Phe Leu Ser  
 35 40 45

Ser Leu Pro Leu His Glu Tyr Pro Pro Ala Phe Pro Leu Gly Ala Asp  
 50 55 60

Ile Gln Gly Leu Asp Leu Phe Ser Phe Leu Gln Thr Glu Ser Gln His  
 65 70 75 80

Tyr Xaa Pro Ser Val Ile Thr Ser Leu Asp Glu Gln Asp Xaa Leu Gly  
 85 90 95

His Phe Phe Gln Xaa Arg Gly Thr Pro Xaa His Phe Leu Gly Pro Leu  
 100 105 110

Ala Pro Thr Leu Gly Ser Ser His Xaa Ser Ala Thr Pro Ala Pro Xaa  
 115 120 125

228844066.ST25.txt

Pro Gly Arg Val Ser Ser Ile Val Ala Pro Gly Xaa Xaa Leu Arg Glu  
130 135 140

Gly His Gly Gly Pro Leu Pro Ser Gly Pro Ser Leu Thr Gly Cys Arg  
145 150 155 160

Ser Asp Ile Xaa Ser Leu Asp  
165

<210> 12  
<211> 26  
<212> PRT  
<213> Synthetic construct

<400> 12

Ala Thr Gly Ala Ala Gly Ala Thr Cys Ala Ala Ala Gly Ala Gly Cys  
1 5 10 15

Thr Thr Thr Ala Cys Cys Gly Ala Cys Gly  
20 25

<210> 13  
<211> 23  
<212> PRT  
<213> Synthetic construct

<400> 13

Thr Cys Ala Gly Thr Cys Cys Ala Gly Gly Gly Ala Ala Ala Thr Cys  
1 5 10 15

Ala Thr Gly Ala Cys Cys Gly  
20